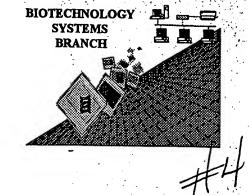
RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following CRF diskette:

Application Serial Number: 09/068,528

Art Unit / Team No.: 1643

Date Processed by STIC: \(\bigcup \frac{0}{26}/98 \)

THE ATTACHED PRINTOUT EXPLAINS THE ERRORS DETECTED.

PLEASE BE SURE TO FORWARD THIS INFORMATION TO THE APPLICANTS BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANTS ALONG WITH A NOTICE TO COMPLY or,
- 2) CALLING APPLICANTS AND FAXING THEM A COPY OF THE PRINTOUT WITH A NOTICE TO COMPLY

THIS WILL INSURE THAT THE NEXT SUBMISSION RECEIVED FROM THEM WILL BE ERROR FREE.

IF YOU HAVE ANY FURTHER QUESTIONS, PLEASE CALL:

ARTI SHAH 703-308-4212

01/008/320 precursor and a sugar as the sole starting materials and of a complex carbohydrate from the sugar nucleotide and a complex carbohydrate precursor. SEQUENCE LISTING SEQ ID NO:1:(2) INFORMATION FOR SER ID NO: 1: Does Not Comply Carrected Diskette Needed Sym -7(1) SEQUENCE CHARACTERISTICSI (A) SEQUENCE LENGTH: 31 (B) SEQUENCE TYPE: nucleic acid (SMANDEDNESS: "Circular," Circular," both MOLECULE TYPE: other nucleic acid, synthetic DNA or tupes on one (i) SEQUENCE DESCRIPTION: SEQ FONO: 1: "un known GGAGAAAGCT TATGGCTGCC ATTAATACGA A SEQ ID NO:2: SEQUENCE LENGTH: 30 (1) GENERAL INFORMATION: (1) APPLICANT: (a) APPLICANT:
(b) TITLE OF INVENTION:
(cii) NUMBER OF SEQUENCES:
(d) ADDRESSEE:
(d) ADDRESSEE:
(e) STREET:
(c) CITY:
(d) STATE:
(e) COUNTRY:
(f) ZIP:
(v) COMPUTER READABLE FORM:
(A) MEDIUM TYPE:
(B) COMPUTER:
(C) OPPRATING SYSTEM:
(D) SOFTWARE: SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded MOLECULAR TYPE: other nucleic acid, synthetic DNA SEQUENCE DESCRIPTION: AACACGGATC CGGATGTTAC TTCTTAATGC SEO ID NO:3: SEQUENCE LENGTH: 28 (D) SOFTWARE:
(vi) CURRENT APPLICATION DATA: SEQUENCE TYPE: nucleic acid (A) APPLICATION NUMBER: (B) FILING DATE: (C) CLASSIFICATION: TOPOLOGY: single-stranded MOLECULAR TYPE: other nucleic acid, synthetic DNA SEQUENCE DESCRIPTION: ATGGAGGATC CTGCTCTGTA TACCGTCT fless 28

follow Significante

format.

also, see 20

sample Jesting aborled in bock for valid 31 28 SEO ID NO:4: SEQUENCE LENGTH: 20 SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded MOLECULAR TYPE: other nucleic acid, synthetic DNA SEQUENCE DESCRIPTION: TGCTGGTCGA CCTGCGCTTG SEQ ID NO:5: SEQUENCE LENGTH: 31 SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded MOLECULAR TYPE: other nucleic acid, synthetic DNA SEQUENCE DESCRIPTION: AAGGAAAGCT TATGACGCAA TTTAATCCCG T SEQ ID NO:6: SEQUENCE LENGTH: 20 SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded MOLECULAR TYPE: other nucleic acid, synthetic DNA SEQUENCE DESCRIPTION: GCAAAGTTAA CAGTCGGTAC 20 SEO ID NO:7: SEQUENCE LENGTH: 31 SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded MOLECULAR TYPE: other nucleic acid, synthetic DNA

09/068,578 SEQUENCE DESCRIPTION: TCAGGAAGCT TATGTTGAAT AATGCTATGA G 31 SEQ ID NO:8: SEQUENCE LENGTH: 27 SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded MOLECULAR TYPE: other nucleic acid, synthetic DNA SEQUENCE DESCRIPTION: TCTCCGGATC CCATGTGACC GGGTTAG 27 SEQ ID NO:9: SEQUENCE LENGTH: 28 SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded MOLECULAR TYPE: other nucleic acid, synthetic DNA SEQUENCE DESCRIPTION: TCTAAATCGA TGCAGACAAA GGACAAAG 28 SEQ ID NO:10: SEQUENCE LENGTH: 27 SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded MOLECULAR TYPE: other nucleic acid, synthetic DNA SEQUENCE DESCRIPTION: TTGCAGGATC CTCGTAGGCC TGATAAG 27 SEQ ID NO:11: SEQUENCE LENGTH: 20 SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded MOLECULAR TYPE: other nucleic acid, synthetic DNA SEQUENCE DESCRIPTION: TGATATCCGC TCCCTTTCCG 20 SEQ ID NO:12: SEQUENCE LENGTH: 26 SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded MOLECULAR TYPE: other nucleic acid, synthetic DNA SEQUENCE DESCRIPTION: ACAGCGGATC CGATGTGTTC GCTGAG 26 SEQ ID NO:13: SEQUENCE LENGTH: 29 SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded MOLECULAR TYPE: other nucleic acid, synthetic DNA SEQUENCE DESCRIPTION: ACAGCAAGCT TTTGACTTTA GCGGAGCAG 29 SEQ ID NO:14: SEQUENCE LENGTH: 29 SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded MOLECULAR TYPE: other nucleic acid, synthetic DNA SEQUENCE DESCRIPTION: GAGTTGGATC CCGATATAAA AGGAAGGAT 29 SEQ ID NO:15:

09/08,528 SEQUENCE LENGTH: 25 SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded MOLECULAR TYPE: other nucleic acid, synthetic DNA SEQUENCE DESCRIPTION: TTTTTAAGCT TCATTTATCA AGAGT 25 SEQ ID NO:16: SEQUENCE LENGTH: 31 SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded MOLECULAR TYPE: other nucleic acid, synthetic DNA SEQUENCE DESCRIPTION: TTTTTGATAT CCCCAATGCT GGGGGTTTTT G 31 SEQ ID NO:17: SEQUENCE LENGTH: 31 SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded MOLECULAR TYPE: other nucleic acid, synthetic DNA SEQUENCE DESCRIPTION: CGTCAAAGCT TAAATGATAT TCGGGGATAA T 31 SEQ ID NO:18: SEQUENCE LENGTH: 25 SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded MOLECULAR TYPE: other nucleic acid, synthetic DNA SEQUENCE DESCRIPTION: AGGGAGGATC CGACATTACT CGTTC 25 SEQ ID NO:19: SEQUENCE LENGTH: 33 SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded MOLECULAR TYPE: other nucleic acid, synthetic DNA SEQUENCE DESCRIPTION: CCGCAAGATC TCGTAAAAAG GGTATCGATA AGC 33 SEQ ID NO:20: SEQUENCE LENGTH: 27 SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded MOLECULAR TYPE: other nucleic acid, synthetic DNA SEQUENCE DESCRIPTION: TTGGGAAGCT TCCGGCAAAT GTGGTTT 27 SEQ ID NO:21: SEQUENCE LENGTH: 25 SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded MOLECULAR TYPE: other nucleic acid, synthetic DNA SEQUENCE DESCRIPTION: ATAAACTCGA GAGAGACAAG CGGAG 25 SEQ ID NO:22:

SEQUENCE LENGTH: 27

SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded

MOLECULAR TYPE: other nucleic acid, synthetic DNA

SEQUENCE DESCRIPTION:

TATTATCGAT GAATTAATAA TTCATAG

27

SEQ ID NO:23:

SEQUENCE LENGTH: 25

SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded

MOLECULAR TYPE: other nucleic acid, synthetic DNA

SEQUENCE DESCRIPTION:

CTCTGGATCC AGTTACGTAT AATAT

25

SEQ ID NO:24:

SEQUENCE LENGTH: 30

SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded

MOLECULAR TYPE: other nucleic acid, synthetic DNA

SEQUENCE DESCRIPTION:

CGGCAAGCTT ATTGTGCCTT TCCAATAAAA

SEQ ID NO:25:

SEQUENCE LENGTH: 28

SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded

MOLECULAR TYPE: other nucleic acid, synthetic DNA

SEQUENCE DESCRIPTION:

ACTTGGATCC CCGTCAATAA ATCTTGCG

28

SEQ ID NO:26:

SEQUENCE LENGTH: 30

SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded

MOLECULAR TYPE: other nucleic acid, synthetic DNA

SEQUENCE DESCRIPTION:

GGTAAAGCTT ATGCAAAACC ACGTTATCAG

SEQ ID NO:27:

SEQUENCE LENGTH: 29

SEQUENCE TYPE: nucleic acid TOPOLOGY: single-stranded

MOLECULAR TYPE: other nucleic acid, synthetic DNA

SEQUENCE DESCRIPTION:

AAACGGATCC TTATTGGAAA GGCACAATA

29

WHÁT IS CLAIMED IS:

1. A process for producing a sugar nucleotide, which

(sel Adoched)

file costains non dequeue Listing